

CRF Errors Corrected by the STIC Systems Branch

Serial Number: 08/486,913

CRF Processing Date: 3/8/96
 Edited by: me
 Verified by: me (STIC staff)

☐ Changed a file from non-ASCII to ASCII

☐ Changed the margins in cases where the sequence text was "wrapped" down to the next line.

☐ Edited a format error in the Current Application Data section, specifically:

ENTERED

☐ Edited the Current Application Data section with the actual current number. The number inputted by the applicant was ☐ the prior application data; or ☐ other _____

☐ Added the mandatory heading and subheadings for "Current Application Data".

☐ Edited the "Number of Sequences" field. The applicant spelled out a number instead of using an integer.

☐ Changed the spelling of a mandatory field (the headings or subheadings), specifically:

☒ Corrected the SEQ ID NO when obviously incorrect. The sequence numbers that were edited were:

Seq. 8

☐ Inserted or corrected a nucleic number at the end of a nucleic line. SEQ ID NO's edited:

☐ Corrected subheading placement. All responses must be on the same line as each subheading. If the applicant placed a response below the subheading, this was moved to its appropriate place.

☐ Inserted colons after headings/subheadings. Headings edited included:

☐ Deleted extra, invalid, headings used by an applicant, specifically:

☐ Deleted: ☐ non-ASCII "garbage" at the beginning/end of files; ☐ secretary initials/filename at end of file;
☐ page numbers throughout text; ☐ other invalid text, such as _____

☐ Inserted mandatory headings, specifically: _____

☐ Corrected an obvious error in the response, specifically: _____

☐ Edited identifiers where upper case is used but lower case is required, or vice versa.

☐ Corrected an error in the Number of Sequences field, specifically: _____

☒ A "Hard Page Break" code was inserted by the applicant. All occurrences had to be deleted. @ Seq. 1

☐ Deleted **ending** stop codon in amino acid sequences and adjusted the "(A)Length:" field accordingly (error due to a PatentIn bug). Sequences corrected: _____

☐ Other: _____

*Examiner: The above corrections must be communicated to the applicant in the first Office Action. DO NOT send a copy of this form.

3/1/95

RAW SEQUENCE LISTING
PATENT APPLICATION US/08/486,913DATE: 03/08/96
TIME: 16:38:10

INPUT SET: S9193.raw

This Raw Listing contains the General
Information Section and up to the first 5 pages.

SEQUENCE LISTING

ENTERED

1
2
3 (1) General Information:
4
5 (i) APPLICANT: Andrew C. Hiatt
6
7 Floyd Rose
8
9 (ii) TITLE OF INVENTION: 3' PROTECTED NUCLEOTIDES
10 FOR ENZYME CATALYZED
11 TEMPLATE-INDEPENDENT CREATION
12 OF PHOSPHODIESTER
13 BONDS
14
15 (iii) NUMBER OF SEQUENCES: 8
16
17 (iv) CORRESPONDENCE ADDRESS:
18
19 (A) ADDRESSEE: Lyon & Lyon
20 (B) STREET: 633 West Fifth Street
21 Suite 4700
22 (C) CITY: Los Angeles
23 (D) STATE: California
24 (E) COUNTRY: U.S.A.
25 (F) ZIP: 90071
26
27 (v) COMPUTER READABLE FORM:
28
29 (A) MEDIUM TYPE: 3.5" Diskette, 1.44 Mb
30 storage
31 (B) COMPUTER: IBM Compatible
32 (C) OPERATING SYSTEM: IBM P.C. DOS 5.0
33 (D) SOFTWARE: Word Perfect 5.1
34
35 (vi) CURRENT APPLICATION DATA:
36
37 (A) APPLICATION NUMBER:
38 (B) FILING DATE:
39 (C) CLASSIFICATION:
40
41 (vii) PRIOR APPLICATION DATA:
42 Prior applications total,
43 including application
44 described below:
45
46 (viii) ATTORNEY/AGENT INFORMATION:

RAW SEQUENCE LISTING
PATENT APPLICATION US/08/486,913DATE: 03/08/96
TIME: 16:38:12

INPUT SET: S9193.raw

47 (A) NAME: Guise, Jeffrey W.
48 (B) REGISTRATION NUMBER: 34,613
49 (C) REFERENCE/DOCKET NUMBER: 207/145
50
51 (ix) TELECOMMUNICATION INFORMATION:
52
53 (A) TELEPHONE: (619) 552-8400
54 (B) TELEFAX: (619) 552-0159
55 (C) TELEX: 67-3510
56
57 (2) INFORMATION FOR SEQ ID NO: 1:
58
59 (i) SEQUENCE CHARACTERISTICS:
60
61 (A) LENGTH: 12
62 (B) TYPE: nucleic acid
63 (C) STRANDEDNESS: single
64 (D) TOPOLOGY: linear
65
66 (ix) FEATURE:
67
68 (D) OTHER INFORMATION: base number 12 is m7g
69
70
71 (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 1:
72
73 CCCCCCCCC CG 12
74
75
76
77 (2) INFORMATION FOR SEQ ID NO: 2:
78
79 (i) SEQUENCE CHARACTERISTICS:
80
81 (A) LENGTH: 19
82 (B) TYPE: nucleic acid
83 (C) STRANDEDNESS: single
84 (D) TOPOLOGY: linear
85
86 (ii) SEQUENCE DESCRIPTION: SEQ ID NO: 2:
87
88 CCCCCCCCC CCCCCTGCA 19
89
90
91
92 (2) INFORMATION FOR SEQ ID NO: 3:
93
94 (i) SEQUENCE CHARACTERISTICS:
95
96 (A) LENGTH: 20
97 (B) TYPE: nucleic acid
98 (C) STRANDEDNESS: single
99 (D) TOPOLOGY: linear

RAW SEQUENCE LISTING
PATENT APPLICATION US/08/486,913DATE: 03/08/96
TIME: 16:38:14

INPUT SET: S9193.raw

100
101 (ii) SEQUENCE DESCRIPTION: SEQ ID NO: 3:
102
103 CTGCAGGGGG GGGGGGGGGG 20
104
105
106 (2) INFORMATION FOR SEQ ID NO: 4:
107
108 (i) SEQUENCE CHARACTERISTICS:
109
110 (A) LENGTH: 9
111 (B) TYPE: nucleic acid
112 (C) STRANDEDNESS: single
113 (D) TOPOLOGY: linear
114
115 (ii) SEQUENCE DESCRIPTION: SEQ ID NO: 4:
116
117 CCCCCCCC 9
118
119
120 (2) INFORMATION FOR SEQ ID NO: 5:
121
122 (i) SEQUENCE CHARACTERISTICS:
123
124 (A) LENGTH: 9
125 (B) TYPE: nucleic acid
126 (C) STRANDEDNESS: single
127 (D) TOPOLOGY: linear
128
129 (ii) SEQUENCE DESCRIPTION: SEQ ID NO: 5:
130
131 GGGGGGGGG 9
132
133
134 (2) INFORMATION FOR SEQ ID NO: 6:
135
136 (i) SEQUENCE CHARACTERISTICS:
137
138 (A) LENGTH: 5
139 (B) TYPE: nucleic acid
140 (C) STRANDEDNESS: single
141 (D) TOPOLOGY: linear
142
143 (ii) SEQUENCE DESCRIPTION: SEQ ID NO: 6:
144
145 CTGCA 5
146
147 (2) INFORMATION FOR SEQ ID NO: 7:
148
149 (i) SEQUENCE CHARACTERISTICS:
150
151 (A) LENGTH: 9
152 (B) TYPE: nucleic acid

RAW SEQUENCE LISTING
PATENT APPLICATION US/08/486,913DATE: 03/08/96
TIME: 16:38:17

INPUT SET: S9193.raw

153 (C) STRANDEDNESS: single
154 (D) TOPOLOGY: linear
155

156 (ii) SEQUENCE DESCRIPTION: SEQ ID NO: 7:
157

158 CTGCATGCA
159
160

9

161 (2) INFORMATION FOR SEQ ID NO: 8:
162

163 (i) SEQUENCE CHARACTERISTICS:
164

165 (A) LENGTH: 10
166 (B) TYPE: nucleic acid
167 (C) STRANDEDNESS: single
168 (D) TOPOLOGY: linear
169

170 (ii) SEQUENCE DESCRIPTION: SEQ ID NO: 8:
171

172 CTGCATGCAG
173
174
175
176
177

10

RAW SEQUENCE LISTING
PATENT APPLICATION US/08/486,913DATE: 03/08/96
TIME: 16:38:18

INPUT SET: S9193.raw

***** PREVIOUSLY ERRORED SEQUENCES - EDITED *****

161 (2) INFORMATION FOR SEQ ID NO: 8:

162

163 (i) SEQUENCE CHARACTERISTICS:

164

165 (A) LENGTH: 10

166 (B) TYPE: nucleic acid

167 (C) STRANDEDNESS: single

168 (D) TOPOLOGY: linear

169

170 (ii) SEQUENCE DESCRIPTION: SEQ ID NO: 8:

171

172 CTGCATGCAG

10

173

174

175

176

177

PAGE: 1

SEQUENCE VERIFICATION REPORT
PATENT APPLICATION US/08/486,913

DATE: 03/08/96
TIME: 16:38:19

INPUT SET: S9193.raw

Line

Error

Original Text

RAW SEQUENCE LISTING
PATENT APPLICATION US/08/486,913DATE: 03/08/96
TIME: 09:50:50

INPUT SET: S9193.raw

This Raw Listing contains the General
Information Section and those Sequences
containing ERRORS.

SEQUENCE LISTING

(1) General Information:

(i) APPLICANT: Andrew C. Hiatt

Floyd Rose

**Does Not Comply
Corrected Diskette Needed**(ii) TITLE OF INVENTION: 3' PROTECTED NUCLEOTIDES
FOR ENZYME CATALYZED
TEMPLATE-INDEPENDENT CREATION
OF PHOSPHODIESTER
BONDS

(iii) NUMBER OF SEQUENCES: 8

(iv) CORRESPONDENCE ADDRESS:

(A) ADDRESSEE: Lyon & Lyon
(B) STREET: 633 West Fifth Street
Suite 4700
(C) CITY: Los Angeles
(D) STATE: California
(E) COUNTRY: U.S.A.
(F) ZIP: 90071

(v) COMPUTER READABLE FORM:

(A) MEDIUM TYPE: 3.5" Diskette, 1.44 Mb
storage
(B) COMPUTER: IBM Compatible
(C) OPERATING SYSTEM: IBM P.C. DOS 5.0
(D) SOFTWARE: Word Perfect 5.1

(vi) CURRENT APPLICATION DATA:

(A) APPLICATION NUMBER:
(B) FILING DATE:
(C) CLASSIFICATION:(vii) PRIOR APPLICATION DATA:
Prior applications total,
including application
described below:

RAW SEQUENCE LISTING
PATENT APPLICATION US/08/486,913DATE: 03/08/96
TIME: 09:50:53

INPUT SET: S9193.raw

46 (viii) ATTORNEY/AGENT INFORMATION:
47 (A) NAME: Guise, Jeffrey W.
48 (B) REGISTRATION NUMBER: 34,613
49 (C) REFERENCE/DOCKET NUMBER: 207/145
50
51 (ix) TELECOMMUNICATION INFORMATION:
52
53 (A) TELEPHONE: (619) 552-8400
54 (B) TELEFAX: (619) 552-0159
55 (C) TELEX: 67-3510(2) INFORMATION FOR SEQ ID NO:
56
--> 57 (i) SEQUENCE CHARACTERISTICS:
58
--> 59 (A) LENGTH: 12
--> 60 (B) TYPE: nucleic acid
--> 61 (C) STRANDEDNESS: single
--> 62 (D) TOPOLOGY: linear
63
--> 64 (ix) Features:
65
--> 66 (D) Other Information: base number 12 is m7g
67
68
--> 69 (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 1:
70
71 CCCCCCCCCC CG 12
72
73
74

fix format

ERRORED SEQUENCES FOLLOW:

159 (2) INFORMATION FOR SEQ ID NO: 8:
160
161 (i) SEQUENCE CHARACTERISTICS:
162
163 (A) LENGTH: 10
164 (B) TYPE: nucleic acid
165 (C) STRANDEDNESS: single
166 (D) TOPOLOGY: linear
167
--> 168 (ii) SEQUENCE DESCRIPTION: SEQ ID NO: 10:
169
170 CTGCATGCAG 10
171
172
173
174
175

8

SEQUENCE VERIFICATION REPORT
PATENT APPLICATION US/08/486,913DATE: 03/08/96
TIME: 09:50:57**INPUT SET: S9193.raw**

Line	Error	Original Text
15	Number of Sequences (8) Doesn't Equal Actual Count (7)	(iii) NUMBER OF SEQUENCES: 8
57	Unknown or Misplaced Identifier	(i) SEQUENCE CHARACTERISTICS:
59	Unknown or Misplaced Identifier	(A) LENGTH: 12
60	Unknown or Misplaced Identifier	(B) TYPE: nucleic acid
61	Unknown or Misplaced Identifier	(C) STRANDEDNESS: single
62	Unknown or Misplaced Identifier	(D) TOPOLOGY: linear
64	Unknown or Misplaced Identifier	(ix) Features:
66	Unknown or Misplaced Identifier	(D) Other Information: base number 12 is m7g
69	Unknown or Misplaced Identifier	(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 1:
168	Wrong Sequence Number	(ii) SEQUENCE DESCRIPTION: SEQ ID NO: 10: